AMENDMENTS TO THE SPECIFICATION

Please replace the paragraph appearing at page 2, lines 4-16, with the following amended paragraph:

In particular, as to the bovine leukemia virus (BLV), which has the gene pX that regulates virus proliferation in the same manner as the human immunodeficiency virus (HIV) and is a retrovirus most related to HTLV-I, a research group in the United States has reported its relationship with the bovine MHC (BoLA) haplotypes mainly focusing disease resistance; however, its relationship with possibility of onset of the leukemia has not been reported. The ratio of cattle infected by this virus (infection rate in Japan) is 10 to 20%, and 1 to 2% of the infected cattle develops extremely malignant endemic bovine leukemia to die after a long latent period of 10 to 15 years. Therefore, economic loss of stockbreeders caused by the virus is very serious. If a possibility of the onset of cattle after BLV infection can be evaluated by the analysis of bovine MHC (BoLA) haplotypes, it becomes possible to preliminarily select disease resistant cattle for bleeding breeding beforehand, and it is expected that extremely safe cattle breeding can be continued.